

AMENDMENTS TO THE CLAIMS

Please amend the claims as follows:

Claim 1 (Currently Amended): A recombinant *E. coli* host cell which is genetically modified for synthesis of a polyketide,

wherein said modification comprises

incorporation of a propionyl CoA carboxylase (pcc) expression system comprising the pccB and accA2 genes from S. coelicolor wherein said pcc expression system produces an enzyme capable of synthesizing 2S-methylmalonyl CoA,

incorporation of at least one expression system for a modular polyketide synthase (PKS), and

incorporation of at least one expression system for a phosphopantetheinyl transferase that phosphopantetheinylates the PKS.

Claims 2-54 (Cancelled)

Claim 55 (Currently Amended): The host cell as in claim 54 1
wherein the host cell further comprises an expression system for biotin ligase which is the *birA* gene from *E. coli*.

Claim 56 (Previously Presented): The host cell as in claim 1
wherein the phosphopantetheinyl transferase expression system comprises the *sfp* gene from *Bacillus subtilis*.

Claim 57 (Cancelled)

Claim 58 (Currently Amended): The host cell as in claim 1
wherein the cell's *prpA-D* operon is disabled deleted or not expressed.

Claim 59 (Previously Presented): The host cell as in claim 1
wherein the PKS is deoxyerythronolide B synthase (DEBS).

Claim 60 (Previously Presented): The host cell of claim 1 wherein the polyketide is
6-deoxyerythronolide B (6-dEB).

Claim 61 (Currently Amended): A recombinant *Streptomyces* host cell which is
genetically modified for enhanced synthesis of a polyketide,
wherein said modification comprises incorporation of ~~an added~~ the matBC gene from
Streptomyces coelicolor or the matBC gene from Rhizobium trifoli wherein the matBC gene is in
addition to endogenous matBC.

Claim 62 (Cancelled)

Claim 63 (Previously Presented): The host cell as in claim 62 61
wherein the modification further comprises incorporation of a the matA gene from
Rhizobium trifoli.

Claim 64 (Previously Presented): The host cell as in claim 61
wherein said modification further comprises incorporation of at least one expression system
for a modular polyketide synthase (PKS).

Claim 65 (Previously Presented): The host cell as in claim 61
wherein the host cell is *Streptomyces coelicolor*.

Claim 66 (Currently Amended): The host cell as in claim 61
wherein the matBC gene is from *Rhizobium trifoli*.

Claim 67 (Currently Amended): The host cell as in claim 64 64
wherein the PKS is DEBS.

Claim 68 (Previously Presented): The cell as in claim 61
wherein the polyketide is 6-dEB.

Claim 69 (Currently Amended): A recombinant *E. coli* host cell which is genetically modified for synthesis of a polyketide,
wherein said modification comprises
incorporation of a matB the matBC gene from *Streptomyces coelicolor* or the matBC gene from Rhizobium trifoli, and
incorporation of at least one expression system for a modular polyketide synthase (PKS), and
incorporation of at least one expression system for a phosphopantetheinyl transferase that phosphopantetheinylates the PKS.

Claim 70 (Cancelled)

Claim 71 (Currently Amended): The host cell as in claim [[70]] 69
wherein the modification further comprises incorporation of a the matA gene from *Rhizobium trifoli*.

Claim 72 (Currently Amended): The host cell as in claim 69
wherein the matBC gene is from *Rhizobium trifoli*.

Claim 73 (Previously Presented): The host cell as in claim 69
wherein the PKS is DEBS.

Claim 74 (Previously Presented): The host cell as in claim 69 wherein the polyketide is 6-dEB.

Claim 75 (Withdrawn): A method to produce a polyketide which method comprises culturing the cells of claim 1 under conditions wherein said polyketide is produced.

Claim 76 (Withdrawn): A method to assess the results of a procedure effecting modification of polyketide synthase genes according to claim 1, resulting in a mixture of said modified genes which method comprises

transfected a culture of *E. coli* of claim 1 with said mixture of modified genes, culturing individual colonies of said transformed *E. coli*, and assessing each colony for polyketide production.

Claim 77 (Currently Amended): The method of claim 75 which further includes providing a substrate, wherein the substrate is of the formula $R_2C(COOH)_2$ $RCH(COOH)_2$ wherein one R is H, methyl or ethyl and the other is H.

Claim 78 (New): A method to produce a polyketide which method comprises culturing the cells of claim 61 under conditions wherein said polyketide is produced.

Claim 79 (New): A method to assess the results of a procedure effecting modification of polyketide synthase genes according to claim 61, resulting in a mixture of said modified genes which method comprises

transfected a culture of *Streptomyces* of claim 61 with said mixture of modified genes, culturing individual colonies of said transformed *Streptomyces*, and assessing each colony for polyketide production.

Claim 80 (New): The method of claim 61 which further includes providing a substrate, wherein the substrate is of the formula $RCH(COOH)_2$ wherein R is H, methyl or ethyl.

Claim 81 (New): A method to produce a polyketide which method comprises culturing the cells of claim 69 under conditions wherein said polyketide is produced.

Claim 82 (New): A method to assess the results of a procedure effecting modification of polyketide synthase genes according to claim 69, resulting in a mixture of said modified genes which method comprises

transfecting a culture of *E. coli* of claim 69 with said mixture of modified genes, culturing individual colonies of said transformed *E. coli*, and assessing each colony for polyketide production.

Claim 83 (New): The method of claim 69 which further includes providing a substrate, wherein the substrate is of the formula RCH(COOH)₂ wherein R is H, methyl or ethyl.